	CF. Priors Corrected by the STIC Sys Branch CRF Processing Dates /// 200 / W
Serial N	Changed a file from non-ASCII to ASCII ENT
LJ.	Changed the margins in cases where the sequence text was want to the next line.
	Edited a format error in the Current Application Data section, specifically:
	Edited the Current Application Data section with the actual current number. The number insutted by the applicant was the prior application data; or other
\Box	Added the mandatory heading and subheadings for "Current Application Data".
	Edited the "Number of Sequences" field. The applicant spelled out a number instead of using an integer.
	Changed the spelling of a mandatory field (the headings or subheadings), specifically:
	Corrected the SEQ ID NO when obviously incorrect. The sequence numbers that were edited were:
	Inserted or corrected a nucleic number at the end of a nucleic line. SEQ ID NO's edited:
	Corrected subheading placement. All responses must be on the same line as each subheading. If the applicant placed a response below the subheading, this was moved to its appropriate place.
	Inserted colons after headings/subheadings. Headings edited included:
	Deleted extra, invalid, headings used by an applicant, specifically:
	Deleted: non-ASCII garbage at the beginning/end of files; secretary initials/filename at end of file; page numbers throughout text; other invalid text, such as
	Inserted mandatory headings, specifically:
	Corrected an obvious error in the response, specifically:
	Edited identifiers where upper case is used but lower case is required, or vice versa.
	Corrected an error in the Number of Sequences field, specifically:
	A "Hard Page Break" code was inserted by the applicant. All occurrences had to be deleted.
	Deleted ending stop codon in amino acid sequences and adjusted the "(A)Length:" field accordingly (error due to a Patentin bug). Sequences corrected:
	Other:
	Tape 20
	A 5
ŁExamin	er: The above corrections must be communicated to the applicant in the first Office 3/1/95 -Action. DO NOT send a copy of this form:

R. Shukla

TECH CENTER ISOURSON

Does Not Comply

Corrected Diskette Needed

RAW SEQUENCE LISTING

PATENT APPLICATION: US/09/101,423B

DATE: 01/11/2001 TIME: 13:19:54

Input Set : A:\Sequence Listing

Output Set: N:\CRF3\01112001\1101423B.raw

3 <110> APPLICANT: Rudland, Philip S.

Barraclough, Roger B.

W--> 5 <120> TITLE OF INVENTION: Metastasis Inducing DNA's

W--> 6 <130> FILE REFERENCE: WPT 0114 PUS

W--> 7 <140> CURRENT APPLICATION NUMBER: US 09/101,423B

8 <141> CURRENT FILING DATE: 1998-11-27

9 <150> PRIOR APPLICATION NUMBER: PCT/GB97/00074

10 <151> PRIOR FILING DATE: 1997-01-10

W--> 11 <160> NUMBER OF SEQ ID: 8

ERRORED SEQUENCES

271 <210> SEQ ID NO: 8

272 <211> LENGTH: 47 base pairs

273 <212> TYPE: DNA

274 <213> ORGANISM: Homo sapiens

W--> 275 <400> SEQUENCE: 8

277 agctataatg eggeegeata tteggeetga teggeegeaa gettgga

W--> 279/M:\clients\W\WPT\0114pusa\2nd revis seq list 12-20-00.wpd E--> 280 kathy 12/20/00

VERIFICATION SUMMARY . DATE: 01/11/2001
PATENT APPLICATION: US/09/101,423B TIME: 13:19:56

Input Set : A:\Sequence Listing Output Set: N:\CRF3\01112001\1101423B.raw

L:5 M:283 W: Missing Blank Line separator, <120> field identifier L:5 M:283 W: Missing Blank Line separator, <130> field identifier L:6 M:283 W: Missing Blank Line separator, <140> field identifier L:7 M:283 W: Missing Blank Line separator, <160> field identifier L:59 M:283 W: Missing Blank Line separator, <400> field identifier L:275 M:283 W: Missing Blank Line separator, <400> field identifier L:279 M:334 W: (2) Invalid Amino Acid in Coding Region, NUMBER OF INVALID KEYS:5 L:280 M:336 W: Invalid Amino Acid Number in Coding Region, SEQ ID:8

L:280 M:254 E: No. of Bases conflict, LENGTH: Input: 0 Counted:54 SEQ:8 L:280 M:320 E: (1) Wrong Nucleic Acid Designator, NUMBER OF INVALID KEYS:6

L:280 M:112 C: (48) String data converted to lower case,

. . . .

L:280 M:252 E: No. of Seq. differs, <211>LENGTH:Input:47 Found:54 SEQ:8